

## UniProtKB/TrEMBL entry O84627

### Entry information

Entry name	O84627_CHLTR
Primary accession number	O84627
Secondary accession numbers	None
Integrated into TrEMBL on	November 1, 1998
Sequence was last modified on	November 1, 1998 (Sequence version 1)
Annotations were last modified on	October 31, 2006 (Entry version 24)

### Name and origin of the protein

Protein name	CHLPN 76kDa Homolog
Synonyms	None
Gene name	OrderedLocusNames: CT_622
From	Chlamydia trachomatis [TaxID: 813] [HAMAP proteome]
Taxonomy	Bacteria; Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia.

### References

[1] NUCLEOTIDE SEQUENCE [LARGE SCALE GENOMIC DNA].

STRAIN=D/UW-3/Cx;

DOI=10.1126/science.282.5389.754; PubMed=9784136

Stephens R.S., Kalman S., Lammel C.J., Fan J., Marathe R., Aravind L., Mitchell W.P., Olinger L., Tatusov R.L., Zhao Q., Koonin E.V., Davis R.W.;

"Genome sequence of an obligate intracellular pathogen of humans: Chlamydia trachomatis.";

Science 282:754-759(1998).

### Comments

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## Cross-references

Sequence databases	
EMBL	AE001273; AAC68226.1; -; Genomic DNA.
PIR	G71490; G71490.
2D gel databases	
PHCI-2DPAGE	O84627; -.
Genome annotation databases	
GenomeReviews	AE001273 GR; CT 622.
KEGG	ctr:CT622; -.
Other	
Implicit links to	CMR; ProDom; HOGENOM; ModBase; UniRef.

## Keywords

Complete proteome.

## Features

None

## Sequence information

Length: 647 AA

Molecular weight: 68526 Da

CRC64: C0D14C2D74473625 [This is a checksum on the sequence]

MESGPESVSS NQSSMNPIN GQIASNSETK ESTKESEASP SASSSVSSWS FLSSAKHALI  
10 20 30 40 50 60

<http://www.expasy.org/cgi-bin/niceprot.pl/printable?ac=O84627>